

**AMENDMENTS TO THE CLAIMS:**

*Please amend claims as follows:*

1 – 25. (Cancelled).

26. (Currently amended) An ~~[[The]]~~ antagonist of a ligand for binding integrins ~~claim 25~~, in which said antagonist consists essentially of amino acid residues selected from the group consisting of:

i) three, four, five, six, seven, eight, or nine amino acid residues of ~~[[said]]~~ strand A (SEQ ID NO: 2) of domain 1 of ICAM-4 (SEQ ID NO: 1); and

ii) three, four, six, seven, eight or nine amino acid residues of strand G (SEQ ID NO: 3) of domain 1, ~~C, F or G strands or said CE loop~~ of ICAM-4 (SEQ ID NO: 1).

27. (Currently amended) The antagonist of claim ~~[[25]]~~ 26, in which said antagonist consists of three, four, five, six, seven, eight or nine amino acid residues of strand A of defined by ICAM-4 strand A and includes amino acid residues F18, W19 and V20 of ICAM-4.

28. (Currently amended) The antagonist of claim 27, in which said antagonist consists of ~~[[an]]~~ a peptide having the amino acid sequence as defined in SEQ ID NO: 9.

29. (Canceled).

30. (Withdrawn) The antagonist of claim 25, in which said antagonist defined by ICAM-4 strand F includes amino acid residues T91, W93 and R97 of ICAM-4.

31. (Withdrawn) The antagonist of claim 30, in which said antagonist or its active site has or consists essentially of an amino acid sequence as defined in SEQ ID NO: 11.

32. (Withdrawn) The antagonist of claim 25, in which said antagonist defined by ICAM-4 strand G includes amino acid residues R92, A94, T95, S96 and R97 of ICAM-4.

33. (Withdrawn) The antagonist of claim 32, in which said antagonist or its active site has or consists essentially of an amino acid sequence as defined in SEQ ID NO: 10.

34. (Withdrawn) The antagonist of claim 25, in which said antagonist defined by ICAM-4 CE loop includes amino acid residues E151 and T154 of ICAM-4.

35. (Withdrawn) An antagonist of a ligand for an epitope or footprint domain for binding integrins, wherein said antagonist is selected from the group consisting of

(I) a low molecular weight compound which binds to the epitope and/or footprint domain to reduce adhesion between the epitope and/or footprint domain and its ligands;

(II) a peptide comprising amino acid residues of the A, C, F, or G strands of the CE loop of ICAM-4; and

(III) a member selected from the group of peptides, drugs and antibodies that binds an ICAM-4 ligand so as to reduce adhesion of the ligand to the epitope and/or footprint domain;

wherein said epitope comprises a member selected from the group consisting of:

(A) strands A and G of domain 1 of ICAM-4 (SEQ ID NO: 1), wherein said A strand (SEQ ID NO: 2) is defined by amino acid residues 17 to 27 of ICAM-4 and said G strand (SEQ ID NO: 3) is defined by amino acid residues 90 to 100 of ICAM-4,

(B) strands F and G of domain 1 of ICAM-4, wherein said F strand (SEQ ID NO: 4) is defined by amino acid residues 77 to 87 of ICAM-4,

(C) strands A and G of domain 1 of ICAM-4 and further defined by amino acid residues W66 on strand E of domain 1 of ICAM-4 and K118 on strand B of domain 2 of ICAM-4, wherein said E strand is defined by amino acid residues 65 to 75 of ICAM-4 (SEQ ID NO 1) and said B strand (SEQ ID NO: 6) is defined by amino acid residues 116 to 126 of ICAM-4, and

(D) strands F and G of domain 1 of ICAM-4 and further defined by amino acid residues W66 on strand E of domain 1 of ICAM-4 and K118 on strand B of domain 2 of ICAM-4 ;

and wherein said footprint domain comprises a first epitope selected from the group listed as (A)-(D) above and a second epitope comprising strand C and strand F of domain 1 of ICAM-4 and a CE loop of domain 2 of ICAM-4, wherein said C strand (SEQ ID NO: 7) is defined by amino acid residues 47 to 54 of ICAM-4, said F strand (SEQ ID NO: 4) is defined by amino acid residues 77 to 87 of ICAM-4 and said CE loop (SEQ ID NO: 8) is defined by amino acid residues 150 to 158 of ICAM-4.

36. (Withdrawn) The antagonist of claim 35, wherein said antagonist is a low molecular weight compound which binds to the epitope and/or footprint domain to reduce adhesion between the epitope and/or footprint domain and its ligands.

37. (Withdrawn) The antagonist of claim 35, wherein said antagonist is a drug that binds an ICAM-4 ligand so as to reduce adhesion of the ligand to the epitope and/or footprint domain.

**38. (New)** The antagonist of claim 27, wherein the integrin is an  $\alpha$ V integrin.

**39. (New)** The antagonist of claim 26, wherein the antagonist consists of three, four, six, seven, eight or nine amino acid residues of strand G (SEQ ID NO: 3) of domain 1 of ICAM-4 and includes amino acid residues R92, A94, T95, S96 and R97 of ICAM-4.

**40. (New)** The antagonist of claim 39, wherein the antagonist consists of a peptide having the amino acid sequence as defined in SEQ ID NO: 10.

**41. (New)** The antagonist of claim 39, wherein the integrin is an  $\alpha$ V integrin or an  $\alpha$ 4 $\beta$ 1 integrin.

**42. (New)** A method of treating an ICAM-4 mediated disease, wherein the method comprises administering to a patient in need thereof an effective amount of an antagonist according to claim 26.

**43. (New)** A method of treating an ICAM-4 mediated disease, wherein the method comprises administering to a patient in need thereof an effective amount of an antagonist of a ligand for binding integrins, wherein the antagonist comprises amino acid residues selected from the group consisting of:

i) three, four, five, six, seven, eight or nine amino acid residues of strand A (SEQ ID NO: 2) of domain 1 of ICAM-4 (SEQ ID NO: 1), and

ii) three, four, six, seven, eight or nine amino acid residues of strand G (SEQ ID NO: 3) of domain 1 of ICAM-4.